

[illegible]

<120> Process for finding oligonucleotide sequences
for nucleic acid amplification methods

<140>

<160> 9

 $\langle 210 \rangle$ 1

<212> DNA

<213> Red sea bream iridovirus gene

ggticaiggt ctaigagaci icccg99

27

<211> 18

<213> Homo sapiens

actccacat agatcact

18

<211> 22

<213> Homo sapiens

<400> 3
ctaiccatgg ctttagtatg ag 22

<210> 4
<211> 21
<212> DNA
<213> Human DNA sequence from PAC 388M5

<400> 4
agcaccctit caggcagtac c 21

<210> 5
<211> 20
<212> DNA
<213> Helianthus tuberosus sst-1 gene

<400> 5
FAM-tgggtcicga aagiccttgt-TAMRA 20

<210> 6
<211> 25
<212> DNA
<213> Drosophila melanogaster

<400> 6
gctcatgitg cacgiictic gagac 25

<210> 7
<211> 20
<212> DNA
<213> cyprinella galactura

<400> 7
catagitcac tcccctgtga 20

<210> 8
<211> 25
<212> DNA
<213> Bovine viral diarrhea virus

<400> 8
aaagigicta gccatgicit tagta

25

<210> 9
<211> 31
<212> DNA
<213> Helianthus tuberosus sst-1 gene

<400> 9
FAM-gtacctgggt cicgaaagic cttgtggtact-TAMRA

31

2025-04-04 14:44:44